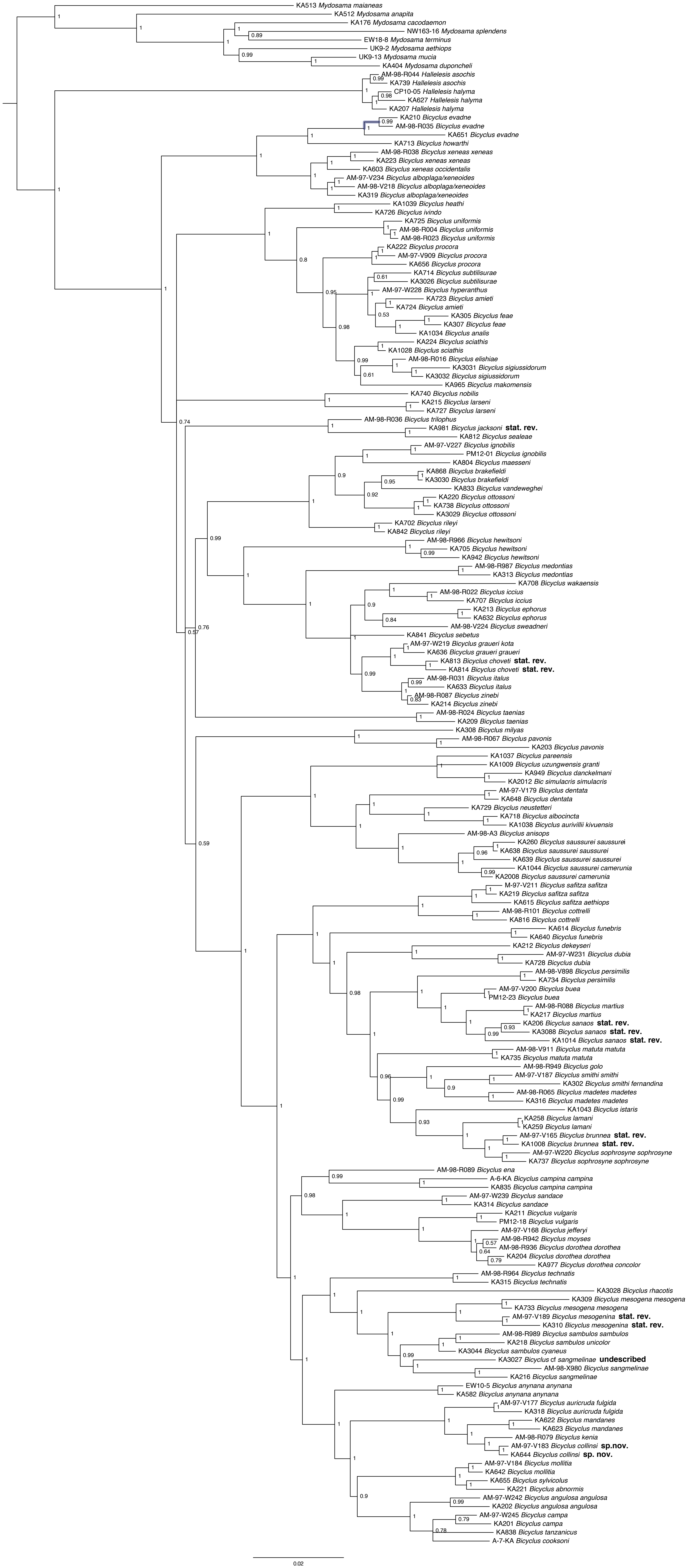


Expanded molecular phylogeny of the genus *Bicyclus* shows the importance of increased sampling for detecting semi-cryptic species and highlights potentials for future studies

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Supplemental data S3: Bayesian Inference tree for the genus *Bicyclus* (with samples from *Mydosama* and *Hallelesis* used as outgroups) constructed by Mr. Bayes 3.2. The numbers at the nodes are the posterior probabilities.



0.02